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## SEQUENCE LISTING

<110> The Scripps Research Institute  
Cheresh, David A.  
Paul, Robert  
Eliceiri, Brian

<120> Method of Treatment of Myocardial  
Infarction

<130> TSRI-651.7

<150> PCT/US03/37653  
<151> 2003-11-18

<150> 10/298,377  
<151> 2002-11-18

<150> 09/538,248  
<151> 2000-03-29

<150> 09/470,881  
<151> 1999-12-22

<150> PCT/US99/11780  
<151> 1999-05-28

<150> 60/087,220  
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cca gcc aac tac gtc cag aag cgg gag ggc gtg aag gcg ggt acc aaa Pro Ala Asn Tyr Val Gln Lys Arg Glu Gly Val Lys Ala Gly Thr Lys	65	70	75
ctc agc ctc atg cct tgg ttc cac ggc aag atc aca cgg gag cag gct Leu Ser Leu Met Pro Trp Phe His Gly Lys Ile Thr Arg Glu Gln Ala	80	85	90
gag cgq ctt ctg tac ccg ccg gag aca ggc ctg ttc ctg gtg cgg gag Glu Arg Leu Leu Tyr Pro Pro Glu Thr Gly Leu Phe Leu Val Arg Glu	95	100	105
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gcc ctg aac atg aag gag ctg aag ctg ctg cag acc atc ggg aag ggg Ala Leu Asn Met Lys Glu Leu Lys Leu Leu Gln Thr Ile Gly Lys Gly	190	195	200
gag ttc gga gac gtg atg ctg ggc gat tac cga ggg aac aaa gtc gcc Glu Phe Gly Asp Val Met Leu Gly Asp Tyr Arg Gly Asn Lys Val Ala	205	210	215
gtc aag tgc att aag aac gac gcc act gcc cag gcc ttc ctg gct gaa Val Lys Cys Ile Lys Asn Asp Ala Thr Ala Gln Ala Phe Leu Ala Glu	225	230	235

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cct cg <sup>g</sup> gtg gag aag ggc tac aag atg gat gcc ccc gac gg <sup>c</sup> tgc ccg Pro Arg Val Glu Lys Gly Tyr Lys Met Asp Ala Pro Asp Gly Cys Pro 400 405 410	1369
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445 450

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Lys Ala Lys Asn Lys Val Gly Arg Glu Gly Ile Ile Pro Ala Asn Tyr  
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Val Gln Lys Arg Glu Gly Val Lys Ala Gly Thr Lys Leu Ser Leu Met  
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Pro Trp Phe His Gly Lys Ile Thr Arg Glu Gln Ala Glu Arg Leu Leu  
85 90 95  
Tyr Pro Pro Glu Thr Gly Leu Phe Leu Val Arg Glu Ser Thr Asn Tyr  
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Pro Gly Asp Tyr Thr Leu Cys Val Ser Cys Asp Gly Lys Val Glu His  
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130 135 140  
Tyr Phe Glu Asn Leu Met Gln Leu Val Glu His Tyr Thr Ser Asp Ala  
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Asp Gly Leu Cys Thr Arg Leu Ile Lys Pro Lys Val Met Glu Gly Thr  
165 170 175  
Val Ala Ala Gln Asp Glu Phe Tyr Arg Ser Gly Trp Ala Leu Asn Met  
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Lys Glu Leu Lys Leu Leu Gln Thr Ile Gly Lys Gly Glu Phe Gly Asp  
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Val Met Leu Gly Asp Tyr Arg Gly Asn Lys Val Ala Val Lys Cys Ile  
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Lys Asn Asp Ala Thr Ala Gln Ala Phe Leu Ala Glu Ala Ser Val Met  
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Asp Cys Leu Leu Lys Phe Ser Leu Asp Val Cys Glu Ala Met Glu Tyr  
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305 310 315 320  
Leu Val Ser Glu Asp Asn Val Ala Lys Val Ser Asp Phe Gly Leu Thr  
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370 375 380  
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cggacggccgg cgcgcgccaga tttgata atg ggc tgc att aaa agt aaa gaa aac 234  
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75	80		85	
aca ggt ggt gtt act ata ttt gtg gcc tta tat gat tat gaa gct aga Thr Gly Gly Val Thr Ile Phe Val Ala Leu Tyr Asp Tyr Glu Ala Arg				522
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Ser Ser Gly Val Thr Pro Phe Gly Gly Ala Ser Ser Ser Phe Ser Val  
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Val Pro Ser Ser Tyr Pro Ala Gly Leu Thr Gly Gly Val Thr Ile Phe  
85 90 95  
Val Ala Leu Tyr Asp Tyr Glu Ala Arg Thr Thr Glu Asp Leu Ser Phe  
100 105 110  
Lys Lys Gly Glu Arg Phe Gln Ile Ile Asn Asn Thr Glu Gly Asp Trp  
115 120 125  
Trp Glu Ala Arg Ser Ile Ala Thr Gly Lys Asn Gly Tyr Ile Pro Ser  
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Asn Tyr Val Ala Pro Ala Asp Ser Ile Gln Ala Glu Glu Trp Tyr Phe  
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Gly Lys Met Gly Arg Lys Asp Ala Glu Arg Leu Leu Leu Asn Pro Gly  
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Ala Tyr Ser Leu Ser Ile Arg Asp Trp Asp Glu Ile Arg Gly Asp Asn  
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Ala Gln Ile Met Lys Lys Leu Arg His Asp Lys Leu Val Pro Leu Tyr  
325 330 335  
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